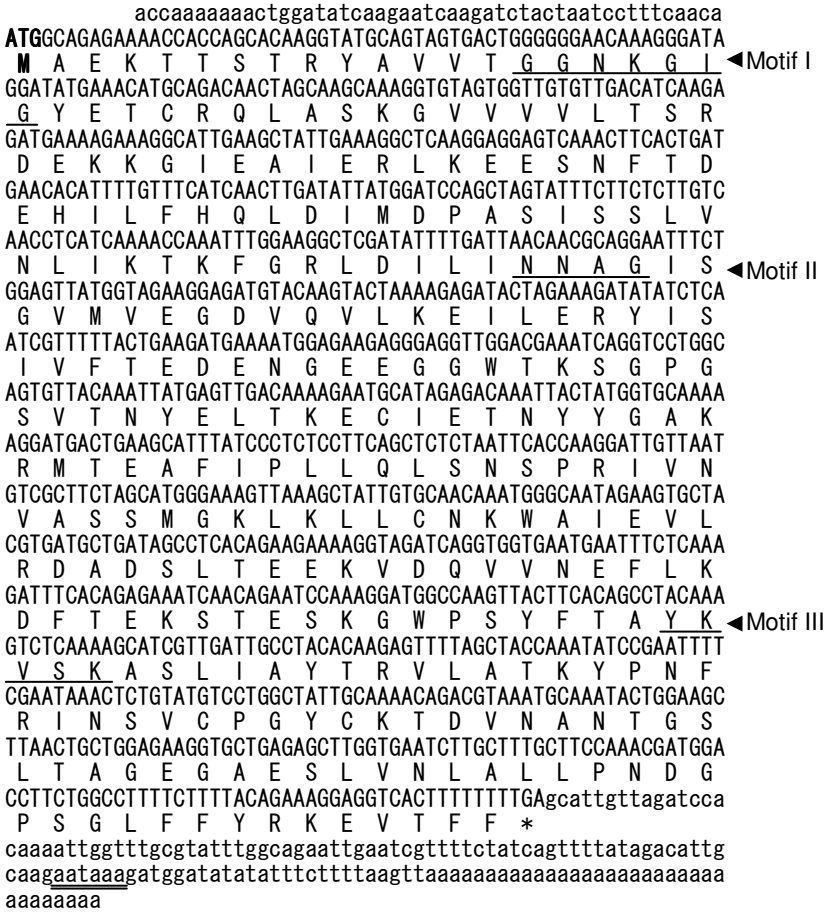

Supplemental Figure S1. Nucleotide sequence and deduced amino acid sequence of the *CaMNR1* cDNA encoding the menthone: (+)-(3s)-neomenthol reductase protein. The deduced amino acid sequence is given below the nucleotide sequence. Translation start site is shown in bold type, termination codon is marked by an asterisk(*) and polyadenylated sequences are double-underlined. The amino acid sequences of conserved domains of SDR are underlined. Binding domain (Motif I: GXXXGXG), the structural domain of undefined function (Motif II: (N/C)NAG), and the active site YXXXK element (Motif III) are shown by upper lines.